

Amendments to the Drawings

The attached sheets of drawings include changes to Figure 2. These sheets, which include Figures 2A-2D, replace the original sheets including Figures 2A-2D. In Figure 2B, nucleotide residues 361-420 and their corresponding amino acid residues have been added.

Attachment: Replacement Sheets

Annotated Sheets Showing Changes

Remarks

Figure 2B has been amended to include nucleotide residues 361-420 and their corresponding amino acid residues. Support for this amendment can be found in the original specification at pages 60-63 (SEQ ID NO:3 and SEQ ID NO:4) and original Figure 2A. Accordingly, these changes are believed to introduce no new matter, and their entry is respectfully requested.

Conclusion

Prompt and favorable consideration of this Preliminary Amendment is respectfully requested. Applicants believe the present application is in condition for allowance. If the Examiner believes, for any reason, that personal communication will expedite prosecution of this application, the Examiner is invited to telephone the undersigned at the number provided.

Respectfully submitted,

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nucleotide residues
361-420 and corresponding
amino acid residues
added

361 CAATAGTGGGGCTGCAGTGCATGAACACCAGCAGCTACCTCAGCAAGACGTTATT
N S E G L Q C M N T S T S Y L S K T L F
421 TGAATTACAGTGCCCTCTCTCTCAAGGCCCCAAACCAGTAACAATCAGTTTGGCCAATCA
E I T V P L S Q G P K P V T I S F A N H
481 CACTTCCTGCCGATGCTCTAACTGGATGTTTACAGACAAGTTCATTCCATTATTAG
T S C R C M S K L D V Y R Q V H S I I R
541 ACGTTCCCTGCCAGCAACACTACCACAGTGTGAGGCGGAGCAAGACCTGCCCCACCAA
R S L P A T L P Q C Q A A N K T C P T N
601 TTACATGTGGAATAATCACATCTGCAGATGCCCTGGCTCAGGAAGATTTATGTTTCCTC
Y M W N N H I C R C L A Q E D F M F S S
661 GGATGCTGGAGATGACTCAACAGATGGATTCCATGACATCTGTGGACCAACAGGAGCT
D A G D D S T D G G F H D I C G P N K E L

FIG. 2B

721 GGATGAAGACCTGTCACTGTCTGTCAGAGCGGGCTTCGGCCTGCCAGCTGTGGACC
-----+-----+-----+-----+-----+-----+
D E E T C C Q C V C R A G L R P A S C G P
781 CCACAAAGAACTAGACAGAACTCATGCCAGTGTGTCTGTAAACAACTCTTCCCCAG
-----+-----+-----+-----+-----+-----+
H K E L D R N S C C Q C V C K N K L F P S
841 CCAATGTGGGCCAACCGAGAATTGATGAAACACATGCCAGTGTGTATGTAAAGAAC
-----+-----+-----+-----+-----+-----+
Q C G A N R E F D E N T C C Q C V C K R T
901 CTGCCCCAGAAATCAACCCCTAAATCCTGGAAATGTGCCCTGTGAATGTACAGAAAGTCC
-----+-----+-----+-----+-----+-----+
C P R N Q P L N P G K C A C E C T E S P
961 ACAGAAATGCTTGTAAAGGAAAGAGTTCCACCACCAACATGCAGCTGTTACAGACG
-----+-----+-----+-----+-----+-----+
Q K C L L K G K K F H H Q T C S C Y R R
1021 GCCATGTACGAACCGCAGAGGCTGTGAGCCAGGATTTTCATATAGTGAAGAAGTGTG
-----+-----+-----+-----+-----+-----+
P C T N R Q K A C E P G F S Y S E E V C

FIG. 2C

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1081 TCGTTGTCCCTTCATATTGGCAAGACCACAAATGAGCTAAGATTGTA CTGTTTCCCA  
-----+-----+-----+-----+-----+-----+  
R C V P S Y W Q R P Q M S  
1141 GTTCATCGATTTTCTATTATGGAAACTGTGTGCCACAGTAGAACTGTCTGTGAACAGA  
-----+-----+-----+-----+-----+-----+  
1201 GAGACCCCTTGTGGTCCATGCTAACAAAGACAAAAGTCTGTCTTTCCCTGAACCATGTGGA  
-----+-----+-----+-----+-----+-----+  
1261 TAACTTTACAGAAATGGACTGGAGCTCATCTGCAAAAGGCCCTCTTGTAAGACTGGTTTT  
-----+-----+-----+-----+-----+-----+  
1321 CTGCCAATGACCAACAGCCAAGATTTTCCCTCTGTGATTTCTTTAAAGAATGACTATA  
-----+-----+-----+-----+-----+-----+  
1381 TAATTATTTCCTCACTAAATAATTGTTTCTGCAATTCATTTTATAGCAACAACAATTGGT  
-----+-----+-----+-----+-----+-----+  
1441 AAACTCACTGTGATCAATAATTTTATATCATGCAGAAATATGTTTAAATAAATGAAAA  
-----+-----+-----+-----+-----+-----+  
1501 TTGTATTATAAAAAAATAAAAAA  
-----+-----+-----+-----+-----+-----+
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FIG. 2D